ELSEVIER

Gene 199 (1997) 311–313

GENE
AN INTERNATIONAL JOURNAL ON GENES AND GENOMES

Author Index Vol. 199 (1997)

Adachi, O., see Izu, H. (199) 203
Alves, G., see Fanning, T. (199) 279
Anderson, A.J., see Kim, Y.C. (199) 219
Aoshima, H., see Izu, H. (199) 203
Arnaud, N., Cheynet, V., Oriol, G., Mandrand, B. and Mallet, F.
Construction and expression of a modular gene encoding bacteriophage T7 RNA polymerase (199) 149
Ashworth, L.K.., see Vinogradova, T. (199) 255
Aubourg, S., Takvorian, A., Chéron, A., Kreis, M. and Lecharny, A.
Structure, organization and putative function of the genes identified within a 23.9-kb fragment from *Arabidopsis thaliana* chromosome IV (199) 241

Bahri, S.M., see Dick, T. (199) 139
Bandman, E., see Chao, T.-H. (199) 265
Beneš, V., see Pečenková, T. (199) 157
Burke, D.T., see Tucker, R.M. (199) 25

Chang, M.C., see Chang, T.M. (199) 225
Chang, T.M., Liu, C.C. and Chang, M.C.
Cloning and sequence analysis of the gene (*eprA1*) encoding an extracellular protease from *Aeromonas hydrophila* (199) 225
Chao, T.-H. and Bandman, E.
Cloning, nucleotide sequence and characterization of a full-length cDNA encoding the myosin heavy chain from adult chicken pectoralis major muscle (199) 265
Chen, D.J., see Ludwig, D.L. (199) 181
Chen, F., see Ludwig, D.L. (199) 181
Chéron, A., see Aubourg, S. (199) 241
Cheynet, V., see Arnaud, N. (199) 149
Chia, W., see Dick, T. (199) 139
Cole, G.T., see Thomas, P.W. (199) 83

Dargis, M., see Handfield, J. (199) 49
Davey, H.W., Ogg, S.L., Husaini, Y., Snell, R.G., Korobko, I.V., Mather, I.H. and Wilkins, R.J.
Structure and sequence of the bovine butyrophilin gene (199) 57
Davies, T.G.E., Theodoulou, F.L., Hallahan, D.L. and Forde, B.G.
Cloning and characterisation of a novel P-glycoprotein homologue from barley (199) 195
Dick, T., Bahri, S.M. and Chia, W.
Drosophila DPP2C1, a novel member of the protein phosphatase 2C (PP2C) family (199) 139

Espín, G., see Mejía-Ruíz, H. (199) 271
Evans, P. and Kemp, J.
Exon/intron structure of the human transferrin receptor gene (199) 123

Fanning, T. and Alves, G.
A family of repetitive DNA sequences in Old World primates (199) 279
Fellous, M., see Veitia, R.A. (199) 63
Flint, A.P.F., see Kaluz, S. (199) 283
Forde, B.G., see Davies, T.G.E. (199) 195
Frelin, L., see Wang, H. (199) 39
Fujii, M., Saijoh, K., Kobayashi, T., Fujii, S., Lee, M.J. and Sumino, K.
Analysis of bovine selenoprotein P-like protein gene and availability of metal responsive element (MRE) located in its promoter (199) 211
Fujii, S., see Fujii, M. (199) 211

Gagnon, L., see Handfield, J. (199) 49
Gomez-Escobar, N., van den Biggelaar, A. and Maizels, R.
A member of the TGF-β receptor gene family in the parasitic nematode *Brugia pahangi* (199) 101
Grzeschik, K.-H., see Vinogradova, T. (199) 255
Guzmán, J., see Mejía-Ruíz, H. (199) 271

Hallahan, D.L., see Davies, T.G.E. (199) 195
Handfield, J., Gagnon, L., Dargis, M. and Huletsky, A.
Sequence of the *ponA* gene and characterization of the penicillin-binding protein 1A of *Pseudomonas aeruginosa* PAO1 (199) 49
Heckmann, S., Schliwa, M. and Kube-Granderath, E.
Primary structure of *Neurospora crassa* γ-tubulin (199) 303
Hernández, R., see Martínez-Calvillo, S. (199) 71
Hinata, K., see Suzuki, G. (199) 133
Hiramatsu, Y., Kato, R., Kawaguchi, S.-i. and Kuramitsu, S.
Cloning and characterization of the *uvrD* gene from an extremely thermophilic bacterium, *Thermus thermophilus* HB8 (199) 77
Huletsky, A., see Handfield, J. (199) 49
Husaini, Y., see Davey, H.W. (199) 57

Ide, N. and Kutsukake, K.
Identification of a novel *Escherichia coli* gene whose expression is dependent on the flagellum-specific sigma factor, FliA, but dispensable for motility development (199) 19
Isogai, A., see Suzuki, G. (199) 133
Izu, H., Kawai, T., Yamada, Y., Aoshima, H., Adachi, O. and Yamada, M.
Characterization of the *gntT* gene encoding a high-affinity gluconate permease in *Escherichia coli* (199) 203

Jin, N., see Kiang, D.T. (199) 165

Kafala, B. and Sasarman, A.
Isolation of the *Staphylococcus aureus hemCDBL* gene cluster coding for early steps in heme biosynthesis (199) 231

Kaluz, S., Kaluzova, M. and Flint, A.P.F.
Sequencing analysis of prion genes from red deer and camel (199) 283
Kaluzova, M., see Kaluz, S. (199) 283
Kato, R., see Hiramatsu, Y. (199) 77
Kawaguchi, S.-i., see Hiramatsu, Y. (199) 77
Kawai, T., see Izu, H. (199) 203
Kaye, F.J., see Otterson, G.A. (199) 287
Kemp, J., see Evans, P. (199) 123
Khil, P., see Vinogradova, T. (199) 255
Kiang, D.T., Jin, N., Tu, Z.-J. and Lin, H.H.
Upstream genomic sequence of the human connexin26 gene (199) 165
Kim, C.H., Oh, Y. and Lee, T.H.
Codon optimization for high-level expression of human erythropoietin (EPO) in mammalian cells (199) 293
Kim, Y.C., Miller, C.D. and Anderson, A.J.
Identification of adjacent genes encoding the major catalase and a bacterioferritin from the plant-beneficial bacterium *Pseudomonas putida* (199) 219
Kirkland, T.N., see Thomas, P.W. (199) 83
Kobayashi, T., see Fujii, M. (199) 211
Korobko, I.V., see Davey, H.W. (199) 57
Kreis, M., see Aubourg, S. (199) 241
Kube-Granderath, E., see Heckmann, S. (199) 303
Kuramitsu, S., see Hiramatsu, Y. (199) 77
Kutsukake, K., see Ide, N. (199) 19

Lavrentyeva, I., see Vinogradova, T. (199) 255
Lebedev, Y., see Vinogradova, T. (199) 255
Lecharny, A., see Aubourg, S. (199) 241
Lee, M.J., see Fujii, M. (199) 211
Lee, T.H., see Kim, C.H. (199) 293
Li, G.C., see Ludwig, D.L. (199) 181
Lin, H.H., see Kiang, D.T. (199) 165
Liu, C.C., see Chang, T.M. (199) 225
López, I., see Martínez-Calvillo, S. (199) 71
Lowe, A.W., see Mandic, R. (199) 173
Ludwig, D.L., Chen, F., Peterson, S.R., Nussenzweig, A., Li, G.C. and Chen, D.J.
Ku80 gene expression is Sp1-dependent and sensitive to CpG methylation within a novel *cis* element (199) 181

Maier, R.J., see Olson, J.W. (199) 93
Maizels, R., see Gomez-Escobar, N. (199) 101
Mallet, F., see Arnaud, N. (199) 149
Mandic, R., Trimble, W.S. and Lowe, A.W.
Tissue-specific alternative RNA splicing of rat vesicle-associated membrane protein-1 (VAMP-1) (199) 173
Mandrand, B., see Arnaud, N. (199) 149
Martínez-Calvillo, S., López, I. and Hernández, R.
pRIBOTEX expression vector: a pTEX derivative for a rapid selection of *Trypanosoma cruzi* transfectants (199) 71
Mather, I.H., see Davey, H.W. (199) 57
McElreavey, K., see Veitia, R.A. (199) 63
Mejía-Ruíz, H., Guzmán, J., Moreno, S., Soberón-Chávez, G. and Espín, G.
The *Azotobacter vinelandii alg8* and *alg44* genes are essential for alginate synthesis and can be transcribed from an *algD*-independent promoter (199) 271
Miller, C.D., see Kim, Y.C. (199) 219
Moreno, S., see Mejía-Ruíz, H. (199) 271

Nelson, B.H., see Soto, J.G. (199) 31
Nussenzweig, A., see Ludwig, D.L. (199) 181

Ocheretina, O. and Scheibe, R.
Cloning and sequence analysis of cDNAs encoding plant cytosolic malate dehydrogenase (199) 145
Ogg, S.L., see Davey, H.W. (199) 57
Oh, Y., see Kim, C.H. (199) 293
Olson, J.W. and Maier, R.J.
The sequences of *hypF*, *hypC* and *hypD* complete the *hyp* gene cluster required for hydrogenase activity in *Bradyrhizobium japonicum* (199) 93
Oriol, G., see Arnaud, N. (199) 149
Otterson, G.A. and Kaye, F.J.
A 'core ATPase', Hsp70-like structure is conserved in human, rat, and *C. elegans* STCH proteins (199) 287

Pačes, J., see Pečenková, T. (199) 157
Pačes, V., see Pečenková, T. (199) 157
Pečenková, T., Beneš, V., Pačes, J., Vlček, Č. and Pačes, V.
Bacteriophage B103: complete DNA sequence of its genome and relationship to other *Bacillus* phages (199) 157
Peterson, S.R., see Ludwig, D.L. (199) 181
Pevsner, J., see Wang, H. (199) 39
Pishko, E.J., see Thomas, P.W. (199) 83

Rayko, E.
Organization, generation and replication of amphimeric genomes: a review (199) 1

Saijoh, K., see Fujii, M. (199) 211
Sasarman, A., see Kafala, B. (199) 231
Scheibe, R., see Ocheretina, O. (199) 145
Schliwa, M., see Heckmann, S. (199) 303
Shevchenko, Y., see Vinogradova, T. (199) 255
Singleton, C.K.
Identification and characterization of the thiamine transporter gene of *Saccharomyces cerevisiae* (199) 111
Snell, R.G., see Davey, H.W. (199) 57
Soberón-Chávez, G., see Mejía-Ruíz, H. (199) 271
Soto, J.G., Nelson, B.H. and Weisblat, D.A.
A leech homolog of *twist*: evidence for its inheritance as a maternal mRNA (199) 31
Sumino, K., see Fujii, M. (199) 211
Suzuki, G., Watanabe, M., Toriyama, K., Isogai, A. and Hinata, K.
Direct cloning of the *Brassica S* locus by using a P1-derived artificial chromosome (PAC) vector (199) 133
Sverdlov, E., see Vinogradova, T. (199) 255

Takvorian, A., see Aubourg, S. (199) 241
Theodoulou, F.L., see Davies, T.G.E. (199) 195
Thomas, P.W., Wyckoff, E.E., Pishko, E.J., Yu, J.-J., Kirkland, T.N. and Cole, G.T.
The *hsp*60 gene of the human pathogenic fungus *Coccidioides immitis* encodes a T-cell reactive protein (199) 83
Toriyama, K., see Suzuki, G. (199) 133
Trimble, W.S., see Mandic, R. (199) 173
Tu, Z.-J., see Kiang, D.T. (199) 165
Tucker, R.M. and Burke, D.T.
Two vectors for the insertion of mammalian selectable genes into yeast artificial chromosome cloned DNA (199) 25

van den Biggelaar, A., see Gomez-Escobar, N. (199) 101
Veitia, R.A., Fellous, M. and McElreavey, K.
Conservation of Y chromosome-specific sequences immediately 5′ to the testis determining gene in primates (199) 63

Vinogradova, T., Volik, S., Lebedev, Y., Shevchenko, Y., Lavrentyeva, I., Khil, P., Grzeschik, K.-H., Ashworth, L.K. and Sverdlov, E.
Positioning of 72 potentially full size LTRs of human endogenous retroviruses HERV-K on the human chromosome 19 map. Occurrences of the LTRs in human gene sites (199) 255
Vlček, Č., see Pečenková, T. (199) 157
Volik, S., see Vinogradova, T. (199) 255

Wang, H., Frelin, L. and Pevsner, J.
Human syntaxin 7: a Pep12p/Vps6p homologue implicated in vesicle trafficking to lysosomes (199) 39
Watanabe, M., see Suzuki, G. (199) 133
Weisblat, D.A., see Soto, J.G. (199) 31
Wilkins, R.J., see Davey, H.W. (199) 57
Wyckoff, E.E., see Thomas, P.W. (199) 83

Yamada, M., see Izu, H. (199) 203
Yamada, Y., see Izu, H. (199) 203
Yu, J.-J., see Thomas, P.W. (199) 83